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   5        Selby, Mark
   6        Houghton, Michael
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5530 ctgatctcct ctgccggaac accgggcatc tccaacttat aagttggaga aataagagaa 12240
5532 tttcagattg agagaatgaa aaaaaaaaac ccttagttca taggtccatt ctcttagcgc 12300
5534 aactacagag aacagggggca caaacaggca aaaaacgggc acaacctcaa tggagtgatg 12360
5536 caacctgcct ggagtaaatg atgacacaag gcaattgacc cacgcatgta tctatctcat 12420
5538 tttcttacac cttctattac cttctgctct ctctgatttg gaaaaagctg aaaaaaaagg 12480
5540 ttgaaaccag ttccctgaaa ttattcccct acttgactaa taagtatata aagacggtag 12540
5542 gtattgattg taattctgta aatctatttc ttaaacttct taaattctac ttttatagtt 12600
5544 agtctttttt ttagtttttaa aacaccaaga acttagtttc gaataaacac acataaacaa 12660
5546 acaagcttac aaaacaaa atg gct gca tat gca gct cag ggc tat aag gtg    12711
5547                      Met Ala Ala Tyr Ala Ala Gln Gly Tyr Lys Val
5548                       1               5                    10
5550 cta gta ctc aac ccc tct gtt gct gca aca ctg ggc ttt ggt gct tac   12759
5551 Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala Tyr
5552         15                  20                   25
5554 atg tcc aag gct cat ggg atc gat cct aac atc agg acc ggg gtg aga   12807
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RAW SEQUENCE LISTING   DATE: 04/30/2001
PATENT APPLICATION: US/09/721,479A   TIME: 08:50:09

Input Set : A:\seqlist.txt
Output Set: N:\CRF3\04302001\I721479A.raw

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5555 Met Ser Lys Ala His Gly Ile Asp Pro Asn Ile Arg Thr Gly Val Arg
5556         30              35              40
5558 aca att acc act ggc agc ccc atc acg tac tcc acc tac ggc aag ttc    12855
5559 Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys Phe
5560         45              50              55
5562 ctt gcc gac ggc ggg tgc tcg ggg ggc gct tat gac ata ata att tgt    12903
5563 Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile Cys
5564 60              65              70              75
5566 gac gag tgc cac tcc acg gat gcc aca tcc atc ttg ggc att ggc act    12951
5567 Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly Thr
5568             80              85              90
5570 gtc ctt gac caa gca gag act gcg ggg gcg aga ctg gtt gtg ctc gcc    12999
5571 Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala
5572             95              100             105
5574 acc gcc acc cct ccg ggc tcc gtc act gtg ccc cat ccc aac atc gag    13047
5575 Thr Ala Thr Pro Pro Gly Ser Val Thr Val Pro His Pro Asn Ile Glu
5576         110             115             120
5578 gag gtt gct ctg tcc acc acc gga gag atc cct ttt tac ggc aag gct    13095
5579 Glu Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys Ala
5580     125             130             135
5582 atc ccc ctc gaa gta atc aag ggg ggg aga cat ctc atc ttc tgt cat    13143
5583 Ile Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys His
5584 140             145             150             155
5586 tca aag aag aag tgc gac gaa ctc gcc gca aag ctg gtc gca ttg ggc    13191
5587 Ser Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu Gly
5588             160             165             170
5590 atc aat gcc gtg gcc tac tac cgc ggt ctt gac gtg tcc gtc atc ccg    13239
5591 Ile Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile Pro
5592         175             180             185
5594 acc agc ggc gat gtt gtc gtc gtg gca acc gat gcc ctc atg acc ggc    13287
5595 Thr Ser Gly Asp Val Val Val Val Ala Thr Asp Ala Leu Met Thr Gly
5596         190             195             200
5598 tat acc ggc gac ttc gac tcg gtg ata gac tgc aat acg tgt gtc acc    13335
5599 Tyr Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Cys Val Thr
5600     205             210             215
5602 cag aca gtc gat ttc agc ctt gac cct acc ttc acc att gag aca atc    13383
5603 Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Ile
5604 220             225             230             235
5606 acg ctc ccc caa gat gct gtc tcc cgc act caa cgt cgg ggc agg act    13431
5607 Thr Leu Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr
5608             240             245             250
5610 ggc agg ggg aag cca ggc atc tac aga ttt gtg gca ccg ggg gag cgc    13479
5611 Gly Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu Arg
5612             255             260             265
5614 ccc tcc ggc atg ttc gac tcg tcc gtc ctc tgt gag tgc tat gac gca    13527
5615 Pro Ser Gly Met Phe Asp Ser Ser Val Leu Cys Glu Cys Tyr Asp Ala
5616         270             275             280
5618 ggc tgt gct tgg tat gag ctc acg ccc gcc gag act aca gtt agg cta    13575
5619 Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala Glu Thr Thr Val Arg Leu
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                 RAW SEQUENCE LISTING              DATE: 04/30/2001
                 PATENT APPLICATION:  US/09/721,479A    TIME: 08:50:09

                 Input Set : A:\seqlist.txt
                 Output Set: N:\CRF3\04302001\I721479A.raw

5620     285                     290                     295
5622 cga gcg tac atg aac acc ccg ggg ctt ccc gtg tgc cag gac cat ctt    13623
5623 Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu
5624 300                     305                     310                     315
5626 gaa ttt tgg gag ggc gtc ttt aca ggc ctc act cat ata gat gcc cac    13671
5627 Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr His Ile Asp Ala His
5628                         320                     325                     330
5630 ttt cta tcc cag aca aag cag agt ggg gag aac ctt cct tac ctg gta    13719
5631 Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu Asn Leu Pro Tyr Leu Val
5632                 335                     340                     345
5634 gcg tac caa gcc acc gtg tgc gct agg gct caa gcc cct ccc cca tcg    13767
5635 Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser
5636             350                     355                     360
5638 tgg gac cag atg tgg aag tgt ttg att cgc ctc aag ccc acc ctc cat    13815
5639 Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr Leu His
5640     365                     370                     375
5642 ggg cca aca ccc ctg cta tac aga ctg ggc gct gtt cag aat gaa atc    13863
5643 Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn Glu Ile
5644 380                     385                     390                     395
5646 acc ctg acg cac cca gtc acc aaa tac atc atg aca tgc atg tcg gcc    13911
5647 Thr Leu Thr His Pro Val Thr Lys Tyr Ile Met Thr Cys Met Ser Ala
5648             400                     405                     410
5650 gac ctg gag gtc gtc acg agc acc tgg gtg ctc gtt ggc ggc gtc ctg    13959
5651 Asp Leu Glu Val Val Thr Ser Thr Trp Val Leu Val Gly Gly Val Leu
5652             415                     420                     425
5654 gct gct ttg gcc gcg tat tgc ctg tca aca ggc tgc gtg gtc ata gtg    14007
5655 Ala Ala Leu Ala Ala Tyr Cys Leu Ser Thr Gly Cys Val Val Ile Val
5656             430                     435                     440
5658 ggc agg gtc gtc ttg tcc ggg aag ccg gca atc ata cct gac agg gaa    14055
5659 Gly Arg Val Val Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu
5660     445                     450                     455
5662 gtc ctc tac cga gag ttc gat gag atg gaa gag tgc tct cag cac tta    14103
5663 Val Leu Tyr Arg Glu Phe Asp Glu Met Glu Glu Cys Ser Gln His Leu
5664 460                     465                     470                     475
5666 ccg tac atc gag caa ggg atg atg ctc gcc gag cag ttc aag cag aag    14151
5667 Pro Tyr Ile Glu Gln Gly Met Met Leu Ala Glu Gln Phe Lys Gln Lys
5668                 480                     485                     490
5670 gcc ctc ggc ctc ctg cag acc gcg tcc cgt cag gca gag gtt atc gcc    14199
5671 Ala Leu Gly Leu Leu Gln Thr Ala Ser Arg Gln Ala Glu Val Ile Ala
5672             495                     500                     505
5674 cct gct gtc cag acc aac tgg caa aaa ctc gag acc ttc tgg gcg aag    14247
5675 Pro Ala Val Gln Thr Asn Trp Gln Lys Leu Glu Thr Phe Trp Ala Lys
5676             510                     515                     520
5678 cat atg tgg aac ttc atc agt ggg ata caa tac ttg gcg ggc ttg tca    14295
5679 His Met Trp Asn Phe Ile Ser Gly Ile Gln Tyr Leu Ala Gly Leu Ser
5680     525                     530                     535
5682 acg ctg cct ggt aac ccc gcc att gct tca ttg atg gct ttt aca gct    14343
5683 Thr Leu Pro Gly Asn Pro Ala Ile Ala Ser Leu Met Ala Phe Thr Ala
5684 540                     545                     550                     555
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5686 gct gtc acc agc cca cta acc act agc caa acc ctc ctc ttc aac ata    14391
5687 Ala Val Thr Ser Pro Leu Thr Thr Ser Gln Thr Leu Leu Phe Asn Ile
5688                 560                 565                 570
5690 ttg ggg ggg tgg gtg gct gcc cag ctc gcc gcc ccc ggt gcc gct act    14439
5691 Leu Gly Gly Trp Val Ala Ala Gln Leu Ala Ala Pro Gly Ala Ala Thr
5692                 575                 580                 585
5694 gcc ttt gtg ggc gct ggc tta gct ggc gcc gcc atc ggc agt gtt gga    14487
5695 Ala Phe Val Gly Ala Gly Leu Ala Gly Ala Ala Ile Gly Ser Val Gly
5696                 590                 595                 600
5698 ctg ggg aag gtc ctc ata gac atc ctt gca ggg tat ggc gcg ggc gtg    14535
5699 Leu Gly Lys Val Leu Ile Asp Ile Leu Ala Gly Tyr Gly Ala Gly Val
5700         605                 610                 615
5702 gcg gga gct ctt gtg gca ttc aag atc atg agc ggt gag gtc ccc tcc    14583
5703 Ala Gly Ala Leu Val Ala Phe Lys Ile Met Ser Gly Glu Val Pro Ser
5704 620                 625                 630                 635
5706 acg gag gac ctg gtc aat cta ctg ccc gcc atc ctc tcg ccc gga gcc    14631
5707 Thr Glu Asp Leu Val Asn Leu Leu Pro Ala Ile Leu Ser Pro Gly Ala
5708                 640                 645                 650
5710 ctc gta gtc ggc gtg gtc tgt gca gca ata ctg cgc cgg cac gtt ggc    14679
5711 Leu Val Val Gly Val Val Cys Ala Ala Ile Leu Arg Arg His Val Gly
5712                 655                 660                 665
5714 ccg ggc gag ggg gca gtg cag tgg atg aac cgg ctg ata gcc ttc gcc    14727
5715 Pro Gly Glu Gly Ala Val Gln Trp Met Asn Arg Leu Ile Ala Phe Ala
5716         670                 675                 680
5718 tcc cgg ggg aac cat gtt tcc ccc acg cac tac gtg ccg gag agc gat    14775
5719 Ser Arg Gly Asn His Val Ser Pro Thr His Tyr Val Pro Glu Ser Asp
5720         685                 690                 695
5722 gca gct gcc cgc gtc act gcc ata ctc agc agc ctc act gta acc cag    14823
5723 Ala Ala Ala Arg Val Thr Ala Ile Leu Ser Ser Leu Thr Val Thr Gln
5724 700                 705                 710                 715
5726 ctc ctg agg cga ctg cac cag tgg ata agc tcg gag tgt acc act cca    14871
5727 Leu Leu Arg Arg Leu His Gln Trp Ile Ser Ser Glu Cys Thr Thr Pro
5728                 720                 725                 730
5730 tgc tcc ggt tcc tgg cta agg gac atc tgg gac tgg ata tgc gag gtg    14919
5731 Cys Ser Gly Ser Trp Leu Arg Asp Ile Trp Asp Trp Ile Cys Glu Val
5732                 735                 740                 745
5734 ttg agc gac ttt aag acc tgg cta aaa gct aag ctc atg cca cag ctg    14967
5735 Leu Ser Asp Phe Lys Thr Trp Leu Lys Ala Lys Leu Met Pro Gln Leu
5736         750                 755                 760
5738 cct ggg atc ccc ttt gtg tcc tgc cag cgc ggg tat aag ggg gtc tgg    15015
5739 Pro Gly Ile Pro Phe Val Ser Cys Gln Arg Gly Tyr Lys Gly Val Trp
5740     765                 770                 775
5742 cga ggg gac ggc atc atg cac act cgc tgc cac tgt gga gct gag atc    15063
5743 Arg Gly Asp Gly Ile Met His Thr Arg Cys His Cys Gly Ala Glu Ile
5744 780                 785                 790                 795
5746 act gga cat gtc aaa aac ggg acg atg agg atc gtc ggt cct agg acc    15111
5747 Thr Gly His Val Lys Asn Gly Thr Met Arg Ile Val Gly Pro Arg Thr
5748                 800                 805                 810
5750 tgc agg aac atg tgg agt ggg acc ttc ccc att aat gcc tac acc acg    15159
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5751 Cys Arg Asn Met Trp Ser Gly Thr Phe Pro Ile Asn Ala Tyr Thr Thr
5752             815                 820                 825
5754 ggc ccc tgt acc ccc ctt cct gcg ccg aac tac acg ttc gcg cta tgg   15207
5755 Gly Pro Cys Thr Pro Leu Pro Ala Pro Asn Tyr Thr Phe Ala Leu Trp
5756             830                 835                 840
5758 agg gtg tct gca gag gaa tac gtg gag ata agg cag gtg ggg gac ttc   15255
5759 Arg Val Ser Ala Glu Glu Tyr Val Glu Ile Arg Gln Val Gly Asp Phe
5760     845                 850                 855
5762 cac tac gtg acg ggt atg act act gac aat ctt aaa tgc ccg tgc cag   15303
5763 His Tyr Val Thr Gly Met Thr Thr Asp Asn Leu Lys Cys Pro Cys Gln
5764 860                 865                 870                 875
5766 gtc cca tcg ccc gaa ttt ttc aca gaa ttg gac ggg gtg cgc cta cat   15351
5767 Val Pro Ser Pro Glu Phe Phe Thr Glu Leu Asp Gly Val Arg Leu His
5768             880                 885                 890
5770 agg ttt gcg ccc ccc tgc aag ccc ttg ctg cgg gag gag gta tca ttc   15399
5771 Arg Phe Ala Pro Pro Cys Lys Pro Leu Leu Arg Glu Glu Val Ser Phe
5772             895                 900                 905
5774 aga gta gga ctc cac gaa tac ccg gta ggg tcg caa tta cct tgc gag   15447
5775 Arg Val Gly Leu His Glu Tyr Pro Val Gly Ser Gln Leu Pro Cys Glu
5776             910                 915                 920
5778 ccc gaa ccg gac gtg gcc gtg ttg acg tcc atg ctc act gat ccc tcc   15495
5779 Pro Glu Pro Asp Val Ala Val Leu Thr Ser Met Leu Thr Asp Pro Ser
5780     925                 930                 935
5782 cat ata aca gca gag gcg gcc ggg cga agg ttg gcg agg gga tca ccc   15543
5783 His Ile Thr Ala Glu Ala Ala Gly Arg Arg Leu Ala Arg Gly Ser Pro
5784 940                 945                 950                 955
5786 ccc tct gtg gcc agc tcc tcg gct agc cag cta tcc gct cca tct ctc   15591
5787 Pro Ser Val Ala Ser Ser Ser Ala Ser Gln Leu Ser Ala Pro Ser Leu
5788             960                 965                 970
5790 aag gca act tgc acc gct aac cat gac tcc cct gat gct gag ctc ata   15639
5791 Lys Ala Thr Cys Thr Ala Asn His Asp Ser Pro Asp Ala Glu Leu Ile
5792             975                 980                 985
5794 gag gcc aac ctc cta tgg agg cag gag atg ggc ggc aac atc acc agg   15687
5795 Glu Ala Asn Leu Leu Trp Arg Gln Glu Met Gly Gly Asn Ile Thr Arg
5796             990                 995                 1000
5798 gtt gag tca gaa aac aaa gtg gtg att ctg gac tcc ttc gat ccg ctt   15735
5799 Val Glu Ser Glu Asn Lys Val Val Ile Leu Asp Ser Phe Asp Pro Leu
5800     1005                1010                1015
5802 gtg gcg gag gag gac gag cgg gag atc tcc gta ccc gca gaa atc ctg   15783
5803 Val Ala Glu Glu Asp Glu Arg Glu Ile Ser Val Pro Ala Glu Ile Leu
5804 1020                1025                1030                1035
5806 cgg aag tct cgg aga ttc gcc cag gcc ctg ccc gtt tgg gcg cgg ccg   15831
5807 Arg Lys Ser Arg Arg Phe Ala Gln Ala Leu Pro Val Trp Ala Arg Pro
5808             1040                1045                1050
5810 gac tat aac ccc ccg cta gtg gag acg tgg aaa aag ccc gac tac gaa   15879
5811 Asp Tyr Asn Pro Pro Leu Val Glu Thr Trp Lys Lys Pro Asp Tyr Glu
5812             1055                1060                1065
5814 cca cct gtg gtc cat ggc tgc ccg ctt cca cct cca aag tcc cct cct   15927
5815 Pro Pro Val Val His Gly Cys Pro Leu Pro Pro Pro Lys Ser Pro Pro
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5816        1070              1075              1080
5818 gtg cct ccg cct cgg aag aag cgg acg gtg gtc ctc act gaa tca acc    15975
5819 Val Pro Pro Pro Arg Lys Lys Arg Thr Val Val Leu Thr Glu Ser Thr
5820    1085              1090              1095
5822 cta tct act gcc ttg gcc gag ctc gcc acc aga agc ttt ggc agc tcc    16023
5823 Leu Ser Thr Ala Leu Ala Glu Leu Ala Thr Arg Ser Phe Gly Ser Ser
5824 1100              1105              1110              1115
5826 tca act tcc ggc att acg ggc gac aat acg aca aca tcc tct gag ccc    16071
5827 Ser Thr Ser Gly Ile Thr Gly Asp Asn Thr Thr Thr Ser Ser Glu Pro
5828              1120              1125              1130
5830 gcc cct tct ggc tgc ccc ccc gac tcc gac gct gag tcc tat tcc tcc    16119
5831 Ala Pro Ser Gly Cys Pro Pro Asp Ser Asp Ala Glu Ser Tyr Ser Ser
5832              1135              1140              1145
5834 atg ccc ccc ctg gag ggg gag cct ggg gat ccg gat ctt agc gac ggg    16167
5835 Met Pro Pro Leu Glu Gly Glu Pro Gly Asp Pro Asp Leu Ser Asp Gly
5836         1150              1155              1160
5838 tca tgg tca acg gtc agt agt gag gcc aac gcg gag gat gtc gtg tgc    16215
5839 Ser Trp Ser Thr Val Ser Ser Glu Ala Asn Ala Glu Asp Val Val Cys
5840    1165              1170              1175
5842 tgc tca atg tct tac tct tgg aca ggc gca ctc gtc acc ccg tgc gcc    16263
5843 Cys Ser Met Ser Tyr Ser Trp Thr Gly Ala Leu Val Thr Pro Cys Ala
5844 1180              1185              1190              1195
5846 gcg gaa gaa cag aaa ctg ccc atc aat gca cta agc aac tcg ttg cta    16311
5847 Ala Glu Glu Gln Lys Leu Pro Ile Asn Ala Leu Ser Asn Ser Leu Leu
5848              1200              1205              1210
5850 cgt cac cac aat ttg gtg tat tcc acc acc tca cgc agt gct tgc caa    16359
5851 Arg His His Asn Leu Val Tyr Ser Thr Thr Ser Arg Ser Ala Cys Gln
5852              1215              1220              1225
5854 agg cag aag aaa gtc aca ttt gac aga ctg caa gtt ctg gac agc cat    16407
5855 Arg Gln Lys Lys Val Thr Phe Asp Arg Leu Gln Val Leu Asp Ser His
5856         1230              1235              1240
5858 tac cag gac gta ctc aag gag gtt aaa gca gcg gcg tca aaa gtg aag    16455
5859 Tyr Gln Asp Val Leu Lys Glu Val Lys Ala Ala Ala Ser Lys Val Lys
5860    1245              1250              1255
5862 gct aac ttg cta tcc gta gag gaa gct tgc agc ctg acg ccc cca cac    16503
5863 Ala Asn Leu Leu Ser Val Glu Glu Ala Cys Ser Leu Thr Pro Pro His
5864 1260              1265              1270              1275
5866 tca gcc aaa tcc aag ttt ggt tat ggg gca aaa gac gtc cgt tgc cat    16551
5867 Ser Ala Lys Ser Lys Phe Gly Tyr Gly Ala Lys Asp Val Arg Cys His
5868              1280              1285              1290
5870 gcc aga aag gcc gta acc cac atc aac tcc gtg tgg aaa gac ctt ctg    16599
5871 Ala Arg Lys Ala Val Thr His Ile Asn Ser Val Trp Lys Asp Leu Leu
5872         1295              1300              1305
5874 gaa gac aat gta aca cca ata gac act acc atc atg gct aag aac gag    16647
5875 Glu Asp Asn Val Thr Pro Ile Asp Thr Thr Ile Met Ala Lys Asn Glu
5876         1310              1315              1320
5878 gtt ttc tgc gtt cag cct gag aag ggg ggt cgt aag cca gct cgt ctc    16695
5879 Val Phe Cys Val Gln Pro Glu Lys Gly Gly Arg Lys Pro Ala Arg Leu
5880    1325              1330              1335
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5882 atc gtg ttc ccc gat ctg ggc gtg cgc gtg tgc gaa aag atg gct ttg    16743
5883 Ile Val Phe Pro Asp Leu Gly Val Arg Val Cys Glu Lys Met Ala Leu
5884 1340          1345          1350          1355
5886 tac gac gtg gtt aca aag ctc ccc ttg gcc gtg atg gga agc tcc tac    16791
5887 Tyr Asp Val Val Thr Lys Leu Pro Leu Ala Val Met Gly Ser Ser Tyr
5888          1360          1365          1370
5890 gga ttc caa tac tca cca gga cag cgg gtt gaa ttc ctc gtg caa gcg    16839
5891 Gly Phe Gln Tyr Ser Pro Gly Gln Arg Val Glu Phe Leu Val Gln Ala
5892          1375          1380          1385
5894 tgg aag tcc aag aaa acc cca atg ggg ttc tcg tat gat acc cgc tgc    16887
5895 Trp Lys Ser Lys Lys Thr Pro Met Gly Phe Ser Tyr Asp Thr Arg Cys
5896          1390          1395          1400
5898 ttt gac tcc aca gtc act gag agc gac atc cgt acg gag gag gca atc    16935
5899 Phe Asp Ser Thr Val Thr Glu Ser Asp Ile Arg Thr Glu Glu Ala Ile
5900     1405          1410          1415
5902 tac caa tgt tgt gac ctc gac ccc caa gcc cgc gtg gcc atc aag tcc    16983
5903 Tyr Gln Cys Cys Asp Leu Asp Pro Gln Ala Arg Val Ala Ile Lys Ser
5904 1420          1425          1430          1435
5906 ctc acc gag agg ctt tat gtt ggg ggc cct ctt acc aat tca agg ggg    17031
5907 Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Leu Thr Asn Ser Arg Gly
5908          1440          1445          1450
5910 gag aac tgc ggc tat cgc agg tgc cgc gcg agc ggc gta ctg aca act    17079
5911 Glu Asn Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr
5912          1455          1460          1465
5914 agc tgt ggt aac acc ctc act tgc tac atc aag gcc cgg gca gcc tgt    17127
5915 Ser Cys Gly Asn Thr Leu Thr Cys Tyr Ile Lys Ala Arg Ala Ala Cys
5916          1470          1475          1480
5918 cga gcc gca ggg ctc cag gac tgc acc atg ctc gtg tgt ggc gac gac    17175
5919 Arg Ala Ala Gly Leu Gln Asp Cys Thr Met Leu Val Cys Gly Asp Asp
5920     1485          1490          1495
5922 tta gtc gtt atc tgt gaa agc gcg ggg gtc cag gag gac gcg gcg agc    17223
5923 Leu Val Val Ile Cys Glu Ser Ala Gly Val Gln Glu Asp Ala Ala Ser
5924 1500          1505          1510          1515
5926 ctg aga gcc ttc acg gag gct atg acc agg tac tcc gcc ccc cct ggg    17271
5927 Leu Arg Ala Phe Thr Glu Ala Met Thr Arg Tyr Ser Ala Pro Pro Gly
5928          1520          1525          1530
5930 gac ccc cca caa cca gaa tac gac ttg gag ctc ata aca tca tgc tcc    17319
5931 Asp Pro Pro Gln Pro Glu Tyr Asp Leu Glu Leu Ile Thr Ser Cys Ser
5932          1535          1540          1545
5934 tcc aac gtg tca gtc gcc cac gac ggc gct gga aag agg gtc tac tac    17367
5935 Ser Asn Val Ser Val Ala His Asp Gly Ala Gly Lys Arg Val Tyr Tyr
5936          1550          1555          1560
5938 ctc acc cgt gac cct aca acc ccc ctc gcg aga gct gcg tgg gag aca    17415
5939 Leu Thr Arg Asp Pro Thr Thr Pro Leu Ala Arg Ala Ala Trp Glu Thr
5940     1565          1570          1575
5942 gca aga cac act cca gtc aat tcc tgg cta ggc aac ata atc atg ttt    17463
5943 Ala Arg His Thr Pro Val Asn Ser Trp Leu Gly Asn Ile Ile Met Phe
5944 1580          1585          1590          1595
5946 gcc ccc aca ctg tgg gcg agg atg ata ctg atg acc cat ttc ttt agc    17511
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5947 Ala Pro Thr Leu Trp Ala Arg Met Ile Leu Met Thr His Phe Phe Ser
5948              1600            1605            1610
5950 gtc ctt ata gcc agg gac cag ctt gaa cag gcc ctc gat tgc gag atc    17559
5951 Val Leu Ile Ala Arg Asp Gln Leu Glu Gln Ala Leu Asp Cys Glu Ile
5952              1615            1620            1625
5954 tac ggg gcc tgc tac tcc ata gaa cca ctg gat cta cct cca atc att    17607
5955 Tyr Gly Ala Cys Tyr Ser Ile Glu Pro Leu Asp Leu Pro Pro Ile Ile
5956         1630            1635            1640
5958 caa aga ctc cat ggc ctc agc gca ttt tca ctc cac agt tac tct cca    17655
5959 Gln Arg Leu His Gly Leu Ser Ala Phe Ser Leu His Ser Tyr Ser Pro
5960     1645            1650            1655
5962 ggt gaa atc aat agg gtg gcc gca tgc ctc aga aaa ctt ggg gta ccg    17703
5963 Gly Glu Ile Asn Arg Val Ala Ala Cys Leu Arg Lys Leu Gly Val Pro
5964 1660            1665            1670            1675
5966 ccc ttg cga gct tgg aga cac cgg gcc cgg agc gtc cgc gct agg ctt    17751
5967 Pro Leu Arg Ala Trp Arg His Arg Ala Arg Ser Val Arg Ala Arg Leu
5968              1680            1685            1690
5970 ctg gcc aga gga ggc agg gct gcc ata tgt ggc aag tac ctc ttc aac    17799
5971 Leu Ala Arg Gly Gly Arg Ala Ala Ile Cys Gly Lys Tyr Leu Phe Asn
5972              1695            1700            1705
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5975 Trp Ala Val Arg Thr Lys Leu Lys Leu Thr Pro Ile Ala Ala Ala Gly
5976         1710            1715            1720
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5979 Gln Leu Asp Leu Ser Gly Trp Phe Thr Ala Gly Tyr Ser Gly Gly Asp
5980     1725            1730            1735
5982 att tat cac agc gtg tct cat gcc cgg ccc cgc tgg atc tgg ttt tgc    17943
5983 Ile Tyr His Ser Val Ser His Ala Arg Pro Arg Trp Ile Trp Phe Cys
5984 1740            1745            1750            1755
5986 cta ctc ctg ctt gct gca ggg gta ggc atc tac ctc ctc ccc aac cga    17991
5987 Leu Leu Leu Leu Ala Ala Gly Val Gly Ile Tyr Leu Leu Pro Asn Arg
5988              1760            1765            1770
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5991 Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
5992              1775            1780            1785
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5995 Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
5996         1790            1795            1800
5998 gga gtt tac ttg ttg ccg cgc agg ggc cct aga ttg ggt gtg cgc gcg    18135
5999 Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
6000     1805            1810            1815
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6003 Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
6004 1820            1825            1830            1835
6006 atc ccc aag gct cgt cgg ccc gag ggc agg acc tgg gct cag ccc ggg    18231
6007 Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Thr Trp Ala Gln Pro Gly
6008              1840            1845            1850
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6011 Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp
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6015 Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
6016      1870           1875           1880
6018 cgg cgt agg tcg cgc aat ttg ggt aag taatagtcga ctttgttccc         18374
6019 Arg Arg Arg Ser Arg Asn Leu Gly Lys
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PATENT APPLICATION: US/09/721,479A   TIME: 08:50:11

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Output Set: N:\CRF3\04302001\I721479A.raw

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